





OIPE

RAW SEQUENCE LISTING

DATE: 04/12/2002

PATENT APPLICATION: US/09/889,936A

TIME: 14:55:51

Input Set : A:\WILLNER5.txt

Output Set: N:\CRF3\04122002\I889936A.raw

ENTERED

3	<110> APPLICANT: WILLNER, Itamar	
4	ESHHAR, Zelig	·
6	<120> TITLE OF INVENTION: DETECTION OF SMALL MOLECULES BY USE (OF A PIEZOELECTRIC
SENSOR		
	<130> FILE REFERENCE: WILLNER=5	
	<140> CURRENT APPLICATION NUMBER: US 09/889,936A	
	<141> CURRENT FILING DATE: 2001-07-25	
	<150> PRIOR APPLICATION NUMBER: PCT/IL00/00048	
	<151> PRIOR FILING DATE: 2000-01-25	
	<150> PRIOR APPLICATION NUMBER: IL 128212	
	<151> PRIOR FILING DATE: 1999-01-25	
	<160> NUMBER OF SEQ ID NOS: 8	
.775	<170> SOFTWARE: PatentIn version 3.1	
23	<210> SEQ ID NO: 1	
24	<211> LENGTH: 37	
	<212> TYPE: DNA	
₁ 26	<213> ORGANISM: Homo sapiens	
[≟28	<400> SEQUENCE: 1	
29	cccgtctaga ggagayatyg twatgaccca gtctcca	37
3	<210> SEQ ID NO: 2	
	<211> LENGTH: 22	
	<212> TYPE: DNA	
<u></u> j=35	<213> ORGANISM: Homo sapiens	
1	<400> SEQUENCE: 2	
38	gtttkatctc gagcttkgts cc	22
41	<210> SEQ ID NO: 3	
42	<211> LENGTH: 17	
	<212> TYPE: DNA	
44	<213> ORGANISM: Homo sapiens	
46	<400> SEQUENCE: 3	
	aggtsmarct kctcgag	17
50	<210> SEQ ID NO: 4	
51	<211> LENGTH: 34	
52	<212> TYPE: DNA	
53	<213> ORGANISM: Homo sapiens	
55	<400> SEQUENCE: 4	
56	tgmrgagacg gtgaccgtrg tyccttggcc ccag	34
59	<210> SEQ ID NO: 5	
60	<211> LENGTH: 408	
61	<212> TYPE: DNA	
62	<213> ORGANISM: Homo sapiens	
64	<220> FEATURE:	
65	<221> NAME/KEY: CDS	
	.000. *********************************	

66 <222> LOCATION: (1)..(408)



RAW SEQUENCE LISTING DATE: 04/12/2002 PATENT APPLICATION: US/09/889,936A TIME: 14:55:51

Input Set : A:\WILLNER5.txt

Output Set: N:\CRF3\04122002\1889936A.raw

	<223					: NOI											
	<400																
	atg																48
72	Met	Arg	Val	Leu	Ile	Leu	Leu	Trp	Leu		Thr	Ala	Phe	Pro		Ile	
73					5				•	10					15		
	ctg																96
76	Leu	Ser	Asp	Val	Gln	Leu	Gln	Glu	Ser	Gly	Pro	Gly	Leu	Val	Lys	Pro	
77				20	•				25					30			
	tcc																144
80	Ser	Gln	Ser	Leu	Ser	Leu	Thr	Cys	Ser	Val	Thr	Gly	Tyr	Ser	Ile	Thr	
81			35					40					45				
83	agt	ggt	tat	gcc	tgg	aac	tgg	atc	cgg	cag	ttt	cca	gga	aac	aaa	ctg	192
	Ser																
85		50	-		-		55		-			60	-		-		
	gag		atσ	aac	tac	ata	aσc	tac	agt.	aat.	ttc	act	agc	tac	aac	cca	240
	Glu																
89 88 <u></u>			1100	011	-1-	70	-	-1-	001	0-1	75			-1-		80	
	tct	ata	242	ant	COL		tat	ttc	act	спа		aca	tcc	aad	aac		288
100 T	Ser	Tan	aya Ama	Cor	N TO	Tla	Cor	Dho	Thr	Ara	Acn	Thr	Cor	Luc	Acn	Gln	200
93	ser	Leu	AIG	ser		TTE	ser	rne	TIII		ASP	1111	per	цуз	95	GIII	
					85					90						+-+	226
	ttc																336
	Phe	Phe	Leu		Leu	Asn	ser	vaı		ser	GIU	Asp	Thr		Thr	Tyr	
≟ 97				100					105					110			
	tac																384
	_	Cys			Trp	Asp	Туг			Thi	тут	: Gly			e Asp	Val	
101			115					120					125)			
	} tgg																408
104	l Trr	Gly	7 Glr	ı Gly	Thr	Thr		. Thr	.								
				-													
105	5	130)				135	5									
	5 3 <21	130)): 6		135	5									
<u></u>	5	130 0> 8) SEQ I	D NC			135	5									
108 109	5 3 <21	130 0> S 1> I) SEQ 1 LENG1	D NO	.36		135	;									
门08 归09 归10	5 3 <21 9 <21	130 0> 5 1> 1 2> 1) SEQ I LENGI TYPE:	D NO H: 1	.36	io sa											
回08 回09 回10 回11	5 3 <21 9 <21 0 <21	130 .0> \$.1> I .2> T) SEQ I LENGI TYPE: ORGAN	D NO CH: 1 PRI	.36 ' Hon	no sa											
108 109 110 113	3 <21 3 <21 9 <21 1 <21 3 <40	130 .0> 5 .1> I .2> 7 .3> 0) SEQ I SENGT TYPE: ORGAN SEQUE	D NOTH: 1 PRI	.36 Hon 6		pier	ıs		ı Phe	e Thr	: Ala	n Phe	e Pro	o Gly	, Ile	
113 115	3 <21 9 <21 0 <21 L <21 8 <40 5 Met	130 .0> 5 .1> I .2> 7 .3> 0) SEQ I SENGT TYPE: ORGAN SEQUE	D NOTH: 1 PRI	.36 Hon 6		pier	ıs		Phe	e Thr	. Ala	n Phe	e Pro	O Gl ₃ 15	, Ile	
115	3 <21 9 <21 0 <21 L <21 8 <40 5 Met	130 .0> 5 .1> I .2> I .3> 0 .0> 5	DEQ I LENGT TYPE: DRGAN SEQUE J Val	D NOTH: 1 PRI IISM: ENCE:	HON 6 1 Ile	e Leu	pier Leu	ıs ı Trp) Leu	10					15		
113 115 116 117	5 <21 3 <21 9 <21 1 <21 2 <40 5 Met 5 1 Leu	130 .0> 5 .1> I .2> I .3> 0 .0> 5	DEQ I LENGT TYPE: DRGAN SEQUE J Val	ID NOTE: 1 PRINTENCE: Leu	HON 6 1 Ile	e Leu	pier Leu	ıs ı Trp) Leu	10					15	y Ile s Pro	
113 115 116 119 120	5	130 .0> 5 .1> I .2> T .3> 0 00> 5 . Arg) SEQ I LENGI TYPE: DRGAN SEQUE J Val	ID NO IH: 1 PRI IISM: ENCE: Leu Val	HON 6 1 Ile 5 Glr	e Leu Leu	pier Leu Glr	ıs Trp	Deu 1 Ser 25	10 Gly	y Pro	Gly	, Leu	va:	15 L Lys	s Pro	
113 115 116 120 123	3 <21 3 <21 3 <21 4 <21 4 <21 5 Met 5 1 Leu 3 Ser	130 .0> 5 .1> I .2> T .3> 0 00> 5 . Arg	DEQ ILENGITYPE: DRGANGEQUE I Val	ID NO IH: 1 PRI IISM: ENCE: Leu Val	HON 6 1 Ile 5 Glr	e Leu Leu	pier Leu Glr	ns Trp Glu	Deu 1 Ser 25	10 Gly	y Pro	Gly	, Leu	va:	15 L Lys		
115 115 115 116 120 123	3 <21 3 <21 0 <21 1 <21 3 <40 5 Met 5 1 Deu	130 .0> 8 .1> 1 .2> 1 .3> 0 .00> 8 . Arg	DEQ ILENGTYPE: DRGAN SEQUE J Val ASP ASP	TD NOTE: 1 PRINCE: ENCE: Leu Val 20 Leu Leu	Hom 6 1 Ile 5 Glr	Leu Leu Leu	pier Leu Glr	ns Trp Glu Cys 40	Leu Ser 25 Ser	10 Gly	Pro	Gly	7 Leu 7 Tyı 45	val 30 Sei	15 L Lys	Pro Thr	
113 115 116 120 123 124	3 <21 3 <21 9 <21 1 <21 5 Met 5 1 1 Leu 1 Ser 4 Ser	130 .0> S .1> I .2> T .3> C .00> S . Arg	DEQ ILENGTYPE: DRGAN SEQUE J Val ASP ASP	TD NOTE: 1 PRINCE: ENCE: Leu Val 20 Leu Leu	Hom 6 1 Ile 5 Glr	Leu Leu Leu	pier Leu Glr Thr	ns Trp Glu Cys 40	Leu Ser 25 Ser	10 Gly	Pro	Gly Gly	7 Leu 7 Tyı 45	val 30 Sei	15 L Lys	s Pro	
108 113 115 116 120 123 124 127	3 <21 3 <21 3 <21 4 <21 5 Met 5 1 6 Leu 7 Ser 8	130 .0> S .1> I .2> T .3> C .00> S . Arc	DEEQ JENGT LENGT LYPE: DRGAN BEQUE J Val ASP ASP 1 Ser 35	TD NOTH: 1 PRINTSM: ENCE: Leu Val 20 Leu Ala	Hom 6 Ile 5 Glr Ser	Leu Leu Leu Asn	pier Leu Glr Thr Trp 55	ns Trp Glu Cys 40 O Ile	D Leu 1 Ser 25 3 Ser 2 Arg	10 Gly Val	Pro Thr	Gly Gly Pro 60	y Lev y Tyr 45 o Gly	Val 30 Sei Asi	15 L Lys c Ile	Pro Thr	
108 113 113 115 116 120 124 127 128 131	3 <21 3 <21 3 <21 4 <21 5 <40 5 Met 5 1 Leu 7 Ser 8 L Glu	130 .0> S .1> I .2> T .3> C .00> S . Arc	DEEQ JENGT LENGT LYPE: DRGAN BEQUE J Val ASP ASP 1 Ser 35	TD NOTH: 1 PRINTSM: ENCE: Leu Val 20 Leu Ala	Hom 6 Ile 5 Glr Ser	Leur Leur Asr	pier Leu Glr Thr Trp 55	ns Trp Glu Cys 40 O Ile	D Leu 1 Ser 25 3 Ser 2 Arg	10 Gly Val	Pro Thr Phe	Gly Gly Pro 60	y Lev y Tyr 45 o Gly	Val 30 Sei Asi	15 L Lys c Ile	Pro Leu Pro	
108 113 113 115 116 120 124 127 128 131	3 <21 3 <21 3 <21 4 <21 5 <40 6 Met 5 1 Leu 7 Ser 8 L Glu 2 65	130 .0> 5 .1> 1 .2> 7 .3> 0 .0> 5 . Arg	DEEQ ILENGT TYPE: DRGAN SEQUE TYPE TYPE: TYPE: TYPE: TYPE TYPE TYPE TYPE TYPE TYPE TYPE TYPE	TD NOTE: 1 PRIVISM: ENCE: Leu 20 Privism: Ala	Hom 6 5 Glr Ser Trp	Leu Leu Leu Asn Tle	Leu Glr Thr Trp 55	Trp Glu Cys 40 O Ile	D Leu 1 Ser 25 3 Ser 2 Arg	10 Gly Val Glr Glr	Pro Thr Phe 75	Gly Gly Pro 60 Thi	y Leu y Tyr 45 o Gly	Val 30 Ser Asi	15 L Lys c Ile n Lys c Asr	Fro Thr Leu Pro 80	
108 113 115 116 120 124 127 128 131 132	3 <21 3 <21 3 <21 4 <21 5 <40 6 Met 5 1 9 Leu 7 Ser 8 1 Glu 2 65 5 Ser	130 .0> 5 .1> 1 .2> 7 .3> 0 .0> 5 . Arg	DEEQ ILENGT TYPE: DRGAN SEQUE TYPE TYPE: TYPE: TYPE: TYPE TYPE TYPE TYPE TYPE TYPE TYPE TYPE	TD NOTE: 1 PRIVISM: ENCE: Leu 20 Privism: Ala	Hom 6 1116 5 Glr Ser Trr	Leu Leu Leu Asn Tle	Leu Glr Thr Trp 55	Trp Glu Cys 40 O Ile	D Leu 1 Ser 25 3 Ser 2 Arg	10 Gly Val Glr Gly Gly Arg	Pro Thr Phe 75	Gly Gly Pro 60 Thi	y Leu y Tyr 45 o Gly	Val 30 Ser Asi	15 L Lys c Ile n Lys c Asi	Pro Leu Pro	
108 110 113 115 116 120 123 124 127 128 131 135	3 <21 3 <21 3 <21 4 <21 5 Met 5 1 6 Leu 7 Ser 8 Glu 2 65 5 Ser	130 .0> S .1> I .2> T .3> C .00> S . Arg . Ser . Glr . Glr . Trr	SEQ ILENGITYPE: DRGAN SEQUE Val ASE TYPE SET TYPE Met	ID NOTH: 1 IPRI IISM: ENCE: Leu Val 20 Leu Ala Gly J Ser	Hom 6 1116 5 Glr 1 Ser 1 Trr 7 Tyr 7 Arg 85	Leu Leu Asm 70	pier Leu Glr Thr Trp 55 Ser	Trp Glu Cys 40 Ile Tyr	D Leu 1 Ser 25 5 Ser 2 Arg 2 Ser 2 Thr	10 Gly Val Glr Glr Gly Are	Pro Thr Phe 75 Asp	Gly Gly Pro 60 Thi	y Lev y Tyr 45 O Gly c Ser	Value Service Tyren Lys	15 L Lys c Ile n Lys c Ass S Ass 95	Fro E Thr Leu Pro 80 Gln	
108 110 113 115 116 120 123 124 127 128 131 135 135	3 <21 3 <21 3 <21 4 <21 5 Met 5 Leu 6 Ser 6 Ser 7 Ser 8 Ser 9 Phe	130 .0> S .1> I .2> T .3> C .00> S . Arg . Ser . Glr . Glr . Trr	SEQ ILENGITYPE: DRGAN SEQUE Val ASE TYPE SET TYPE Met	ID NOTE: 11 PRINTED VAIL 20 Part Alace Gly	HOM 6 1116 5 Glr Ser Trp Tyr Arg 85	Leu Leu Asm 70	pier Leu Glr Thr Trp 55 Ser	Trp Glu Cys 40 Ile Tyr	D Leu 1 Ser 25 3 Ser 2 Arg 5 Ser 4 Thr	10 Gly Val Glr Gly Glr Arc 90 Ser	Pro Thr Phe 75 Asp	Gly Gly Pro 60 Thi	y Lev y Tyr 45 O Gly c Ser	Value Service Tyren Lyse Also	15 L Lys c Ile n Lys c Ass S Ass 95 a Thi	Fro Thr Leu Pro 80	·
108 110 111 115 116 120 123 124 125 131 135 136 136	3 <21 3 <21 3 <21 4 <21 5 Met 5 1 6 Leu 7 Ser 8 Glu 6 Ser 6 Ser 9 Phe	130 10> S 1> I 2> T 3> C 10> S Arg 1 Ser 50 Trr 50 Trr 1 Leu	SEQ I LENGI TYPE: DRGAN SEQUE TYPE:	D NOTE: 1 PRIVISM: PR	HOM 6 Ile 5 Glr Trp Tyr Arg 85 Leu	Leu Leu Asn Tle 70 Tle	pier Glr Thr 55 Ser Ser	Trp Glu Cys 40 Dlle Tyr Phe	D Leu 1 Ser 25 Ser 2 Arg Thr 105	10 Gly Val Glr Gly Arc	Property Pherical Phe	Gly Gly F Gly F From 60 F Thi Thi Asp	y Lev y Tyr 45 o Gly r Ser Ser	Value Service Tyren Lyse Alace 110	15 L Lys C Ile D Lys C Asr 95 A Thr	Fro E Thr Leu Pro 80 Gln	





RAW SEQUENCE LISTING DATE: 04/12/2002 PATENT APPLICATION: US/09/889,936A TIME: 14:55:51

Input Set : A:\WILLNER5.txt

Output Set: N:\CRF3\04122002\1889936A.raw

144			115					120					125					
147	Trp	Gly	Gln	Gly	Thr	Thr	Val	Thr										
148	_	130					135											
151	<210)> SI	EQ II	ONO	: 7													
			ENGTI															
			PE:															
			RGAN		Homo	sar	oiens	5										
			EATUI															
			AME/E		CDS													
			CAT			(34	18)											
			THER			-	-											
			EQUE				-											
						att	ato	acc	cag	tct	cca	tcc	tcc	cta	agt.	ata		48
								Thr										
165		9	011		5	,				10					15			
		αca	σσα	σασ	-	atc	act	atg	age		ааσ	tcc	aσt	caσ		cta		96
168																		,
169	DCI	AIU	OLY	20	פעם	vul	1111	1100	25	Cys	1,5	UCI	001	30	001	DCu		
五71	++>	220	2 a t		22+	722	220	220		++~	aca	+ ~ ~	tan		car	222	1	44
与72 与72	Tou	Aan	Cor	aya	λαη	Cln	Tuc	Agn	Tur	Tou	η C C	Trn	mur	Cln	Cln	Tyc	_	. 7 7
173	пеп	ASII	35	ALG	ASII	GIII	nys	40	TYL	Leu	Ата	ııp	45	GIII	GIII	цуз		
				+			-++		a + a	+	~~~	~+ ·		-++	200	~~+	1	92
175																	1	.92
176	Pro	_	GIII	PTO	PTO	гуѕ		ьeu	тте	тут	GTĀ		Pile	116	ALY	ASP		
177		50	_4_				55					60					_	40
179																	2	40
5		GIY	val	Pro	Asp	_	Pne	Thr	GIĀ	ser	_	ser	СТА	Thr	Asp			
181						70					75					80	_	
183																	2	88
184	Thr	Leu	Thr	IIe		ser	Val	GIn	Ala		Asp	Leu	Ala	Val		Tyr		
185					85					90					95		_	
187																	3	36
188	Cys	Gln	Asn	_	His	Ile	Tyr	Pro		Thr	Phe	Gly	Gly		Thr	Lys		
=189				100					105					110			_	
	_	_	ata														3	48
	Leu	Glu	Ile	Lys			•	•										
193			115															
196	<210)> SI	EQ II	ON C	8													
197	<211	l> LI	ENGTI	i: 11	L6				•									
198	<212	2> T	PE:	PRT														
199	<213	3> OI	RGANI	SM:	Homo	sap	piens	3										
201	<400)> SI	EQUE	ICE:	8													
203	Ser	Arg	Gly	Asp	Ile	Val	Met	Thr	Gln	Ser	Pro	Ser	Ser	Leu	Ser	Val		
204	1	•	_	_	5					10					15			
		Ala	Gly	Glu	Lys	Val	Thr	Met	Ser	Cys	Lys	Ser	Ser	Gln	Ser	Leu		
208			-	20	-				25	_	-			30				
	Leu	Asn	Ser		Asn	Gln	Lys	Asn	Tyr	Leu	Ala	Trp	Tyr	Gln	Gln	Lys		
212			35	-			_	40	-			-	45			_		
	Pro	Gly	Gln	Pro	Pro	Lys	Leu	Leu	Ile	Tyr	Gly	Val	Phe	Ile	Arg	Asp		
216		50				-	55			-	•	60			-	•		





RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/889,936A

DATE: 04/12/2002 TIME: 14:55:51

Input Set : A:\WILLNER5.txt

Output Set: N:\CRF3\04122002\1889936A.raw

219	Ser	Gly	Val	${\tt Pro}$	Asp	Arg	Phe	Thr	Gly	Ser	Gly	Ser	Gly	Thr	Asp	Phe
220	65					70					75					80
223	Thr	Leu	Thr	Ile	Ser	Ser	Val	Gln	Ala	Glu	Asp	Leu	Ala	Val	Tyr	Tyr
224					85	•				90					95	
227	~	01 -	7 ~~	7 ~~	TT 4 -	T1.	m	D	m	ጥኮۍ	Dha	C1	C1 **	C1	mh-	T
221	Cys	GIN	ASII	ASP	HIS	тте	TYL	Pro	TAT	TIIT	Pile	GIY	GTA	GIA	THE	гÃг
227	Cys	GIN	ASII	100	HIS	ire	туг	Pro	105	1111	Pile	СТУ	СТУ	110	THE	гÀг
228	Leu			100	HIS	ire	Tyr	PIO	_	1111	Pile	GIY	GIY	_	THE	гàг

į.





VERIFICATION SUMMARY

PATENT APPLICATION: US/09/889,936A

DATE: 04/12/2002 TIME: 14:55:52

Input Set : A:\WILLNER5.txt
Output Set: N:\CRF3\04122002\I889936A.raw

ı

į <u>-</u>